

SEQUENCE LISTING

<110> Cahoon, Rebecca E.
Falco, Saverio Carl
Pember, Stephen O.

<120> Chorismate Biosynthesis Enzymes

<130> BB-1159-A

<140> 09/743,207

<141> 2001-01-04

<150> PCT/US99/16353

<151> 1999-07-20

<150> 60/093,611

<151> 1998-07-21

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| ggcgcgctcc | cggagtcgcg | ccccacgtcc | ctccggttat | ccgtcggccg | ccgtcggccg | 180 |
| gcctccagcc | tagaggtgaa | ggcatcagga | aatgtgttcg | ggaactactt | ccaggttgca | 240 |
| acctatggcg | aatcccatgg | aggggggtgt | ggttgcgtta | tcagtggctg | cccaccaga | 300 |
| attcctctca | ctgaggcaga | catgcaagta | gaactcgata | gaagacgtcc | gggtcaaagt | 360 |
| agaattacaa | ccccaaagaa | ggagactgat | acatgcaaaa | ttctatcagg | gacacatgat | 420 |
| gggatgacta | ctggtacacc | aattcacgtc | tttgtcccaa | acacagatca | aaggggtggt | 480 |
| gattacagtg | aaatgtctaa | ggcgtacaga | ccatcccatg | cagatgcaac | ctatgacttc | 540 |
| aagtatggag | ttagagctgt | gcagggaggt | ggaagggtcat | cagccagaga | aaccattggc | 600 |
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| ttggcatttg | tttctaaagt | gcaccaagtc | gtacttccag | aagatgcagt | tgattatgag | 720 |
| actgtaacct | tggaaacatat | agagagcaac | atcggttagat | gtcctgatcc | agaatatgca | 780 |
| gagaagatga | ttgctgccat | tgatacggta | cgagtttagag | gagattcaat | tggtggggtc | 840 |
| gtcacatgca | ttgcaagaaa | tgttctctgt | ggtcttggct | ctcctgtttt | tgacaaaactt | 900 |
| gaagctgaac | tggcaaaaagc | catgctttct | cttcctgcaa | gcaagggggt | tgagattggc | 960 |
| agtgggttcg | ctggtacgga | ctttactgga | agtgagcata | atgatgagtt | ctatatggat | 1020 |
| gaggctggaa | atgtgaggac | acgaactaat | cgctcaggcg | gtgttcaggg | agggatatca | 1080 |
| aatggtgaaa | ttatttactt | caaagtggct | tttaagccaa | cagcaactat | cggaaagaag | 1140 |
| caaaaatactg | tgtcaaggga | gcatgaggat | gttgaacttt | tggcaagggg | gcgccatgac | 1200 |
| ccctgtgttg | tccctcgagc | tgttctctatg | gtggaatcca | tggctgcgct | ggtcctgatg | 1260 |
| gaccagctca | tggcgcatat | tgcccagtg | gagatgtttc | cgctgaacct | tgccctacaa | 1320 |
| gagcccattg | gctctgctag | cagtgcacat | gaactgtcac | caaacctatc | ataatgtttg | 1380 |
| tcgtggaaca | tgtcccagct | ttccttctat | cgaaattctg | gtctttgcta | agcagtttgc | 1440 |
| aattcggaa | ccccataaac | cctcgactat | tgtacctaga | gataaagtga | acggatatca | 1500 |
| agatagaaat | gcattaatgt | ttttgtgatg | tgtagtataa | ctgatattta | ccccttttct | 1560 |
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<212> PRT

<213> Zea mays

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| Arg | Phe | Leu | Pro | Arg | Gly | Ile | Gly | Ala | Leu | Pro | Glu | Ser | Ala | Pro | Thr |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Ser | Leu | Arg | Leu | Ser | Val | Gly | Arg | Arg | Arg | Arg | Ala | Ser | Ser | Leu | Glu |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Val | Lys | Ala | Ser | Gly | Asn | Val | Phe | Gly | Asn | Tyr | Phe | Gln | Val | Ala | Thr |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Tyr | Gly | Glu | Ser | His | Gly | Gly | Gly | Val | Gly | Cys | Val | Ile | Ser | Gly | Cys |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| Pro | Pro | Arg | Ile | Pro | Leu | Thr | Glu | Ala | Asp | Met | Gln | Val | Glu | Leu | Asp |
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| Asp | Thr | Cys | Lys | Ile | Leu | Ser | Gly | Thr | His | Asp | Gly | Met | Thr | Thr | Gly |
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| Tyr | Ser | Glu | Met | Ser | Lys | Ala | Tyr | Arg | Pro | Ser | His | Ala | Asp | Ala | Thr |
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| Tyr | Asp | Phe | Lys | Tyr | Gly | Val | Arg | Ala | Val | Gln | Gly | Gly | Gly | Arg | Ser |
| | | | 165 | | | | | | 170 | | | | | 175 | |
| Ser | Ala | Arg | Glu | Thr | Ile | Gly | Arg | Val | Ala | Ala | Gly | Ala | Leu | Ala | Lys |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Lys | Ile | Leu | Lys | Leu | Lys | Ser | Gly | Val | Glu | Ile | Leu | Ala | Phe | Val | Ser |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Lys | Val | His | Gln | Val | Val | Leu | Pro | Glu | Asp | Ala | Val | Asp | Tyr | Glu | Thr |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Val | Thr | Leu | Glu | His | Ile | Glu | Ser | Asn | Ile | Val | Arg | Cys | Pro | Asp | Pro |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Glu | Tyr | Ala | Glu | Lys | Met | Ile | Ala | Ala | Ile | Asp | Thr | Val | Arg | Val | Arg |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Gly | Asp | Ser | Ile | Gly | Gly | Val | Val | Thr | Cys | Ile | Ala | Arg | Asn | Val | Pro |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Arg | Gly | Leu | Gly | Ser | Pro | Val | Phe | Asp | Lys | Leu | Glu | Ala | Glu | Leu | Ala |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Lys | Ala | Met | Leu | Ser | Leu | Pro | Ala | Ser | Lys | Gly | Phe | Glu | Ile | Gly | Ser |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Gly | Phe | Ala | Gly | Thr | Asp | Phe | Thr | Gly | Ser | Glu | His | Asn | Asp | Glu | Phe |

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50 55 60

Val Glu Ala Met Val Ala Leu Val Leu Val Asp Gln Leu Met Ala Gln
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Tyr Ala Gln Cys Asn Leu Phe Pro Val Asn Ser Asp Leu Gln Glu Pro
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Leu Val Pro Ile Leu Arg Pro Glu Glu Ala Leu Leu
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<212> DNA
<213> Triticum aestivum

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gattagtgtg caccaagaga tctgttgatc accgaaaata aaagtttgcg gcgtgaacag 960
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<213> Triticum aestivum

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Pro Glu Asp Ala Val Asp Tyr Glu Thr Leu Thr Leu Asp Gln Ile Glu
35 40 45

Ser Asn Ile Cys Arg Cys Pro Asp Pro Glu Tyr Ala Gln Lys Met Ile
50 55 60

Asp Ala Ile Asp Lys Val Arg Val Asn Gly Asn Ser Ile Gly Gly Val
65 70 75 80

Val Thr Cys Ile Ala Arg Asn Val Pro Arg Gly Leu Gly Ser Pro Val
85 90 95

Phe Asp Lys Leu Glu Ala Leu Leu Ala Lys Ala Met Leu Ser Leu Pro

| | | |
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| Ala Ser Lys Gly Phe Glu Ile Gly Ser Gly Phe Ala Gly Thr Asp Leu 115 120 125 | | |
| Thr Gly Ser Glu His Asn Asp Glu Phe Tyr Met Asp Glu Ala Gly Asn 130 135 140 | | |
| Val Arg Thr Arg Thr Asn Arg Ser Gly Gly Val Gln Gly Gly Ile Ser 145 150 155 160 | | |
| Asn Gly Glu Thr Ile Tyr Phe Lys Val Ala Phe Lys Pro Thr Ala Thr 165 170 175 | | |
| Ile Gly Lys Lys Gln Asn Thr Val Thr Arg Asp His Glu Asp Ile Glu 180 185 190 | | |
| Leu Leu Thr Arg Gly Arg His Asp Pro Cys Val Val Pro Arg Ala Val 195 200 205 | | |
| Pro Met Val Glu Thr Met Ala Ala Leu Val Leu Met Asp Gln Leu Met 210 215 220 | | |
| Ala His Val Ala Gln Cys Glu Met Phe Pro Leu Asn Leu Ala Leu Gln 225 230 235 240 | | |
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<210> 7
 <211> 1626
 <212> DNA
 <213> Zea mays

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| ctcatccaat | ctattccgat | gacgaccgtg | cccaagccac | agcaggtggc | gcactcacgg | 120 |
| gcacggctcg | caccccgcg | gatcgccgcc | ttgctggagt | ttgccccagc | ctcctcctcc | 180 |
| ctccgcttcg | ccgtgcaccg | ctgcgcact | gtcgcctag | aggtgaaggc | atctggaaac | 240 |
| acgtttggaa | actactttca | ggttgcaacc | tatggtgaat | ctcatggggg | tggtggttgt | 300 |
| tgtgttatca | gtggttgcc | acctagaatt | ccactcactg | aggcagacct | acaagttgaa | 360 |
| ctcgatcgaa | gacggcccgg | acagagcaga | ataacctcca | caaggaagga | gactgataca | 420 |
| tgcaaaaattc | tgtcagggac | acatgaaggg | gtgactactg | gaacgccaat | tcttgttatt | 480 |
| gtcccaaaaca | cagatcaaat | aggcagtgat | caccgtgaaa | tagccaatgt | gtaccgacct | 540 |
| tctcatgcag | acgcaactta | tgacttcaag | tacggtgtta | gagctgtaca | gggaggtggg | 600 |
| aggtcctcgg | gcagaaaaac | cgttggaagg | gtggctgcag | gggccctccc | caagaaaatt | 660 |
| cttaagctca | aatgtggatt | agagatcttg | tcgtttgttt | ccaaagtgca | tcaggttgtg | 720 |
| ctcccagaag | acgcggttga | ttatgggtct | gtaactttgg | aacagataga | gagcaacatc | 780 |
| gttagatgtc | ctgatccaga | gtacgcagag | aagatgatag | acgcaatcga | cagagtacga | 840 |
| gttcgagggg | attcgggtcg | tggagtgatc | acatgcgtcg | ctagaaacgt | tcctcgcggg | 900 |
| ctcggttctc | ctgtgttcga | caagctcgaa | tccgaactgg | caaaagctat | gctttctatt | 960 |
| cctgcgagca | acgggttcga | gattggcagc | ggattcgccg | ggaccgactt | gacaggaagt | 1020 |
| gagcataatg | atgagtttta | tatggataag | gctggaagtg | tcaggacacg | gactaatcgc | 1080 |
| tcgggtggtg | tgcagggagg | gatatcgaa | ggttgagattg | tgcacttcaa | agttgctttt | 1140 |
| aagccgacac | catctatcgg | ggtgaaacag | aacaccgtgt | caagggagcg | tcagaacggt | 1200 |
| gagcttctag | caagagggcg | ccatgaccca | tgcgtcgccc | ctcgagctgt | tcctgtggtg | 1260 |
| gaatccatgg | ccgcgttggt | cctcatggac | cagctgatgg | cgcacgtggc | tcagtgcgag | 1320 |
| atgttcgcgc | tcaatactgc | acttcaagaa | ccagttggct | ctttctagca | gaggcagagc | 1380 |

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cgatgaagtg aataaggcac ttggtttcct gtgcatttgt acacgtttca tataatgtaa 1560
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<211> 429
<212> PRT
<213> Zea mays

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Ser Ser Leu Arg Phe Ala Val His Arg Cys Arg Thr Ala Arg Leu Glu
      35              40              45

Val Lys Ala Ser Gly Asn Thr Phe Gly Asn Tyr Phe Gln Val Ala Thr
      50              55              60

Tyr Gly Glu Ser His Gly Gly Gly Val Gly Cys Val Ile Ser Gly Cys
      65              70              75              80

Pro Pro Arg Ile Pro Leu Thr Glu Ala Asp Leu Gln Val Glu Leu Asp
      85              90              95

Arg Arg Arg Pro Gly Gln Ser Arg Ile Thr Ser Thr Arg Lys Glu Thr
      100             105             110

Asp Thr Cys Lys Ile Leu Ser Gly Thr His Glu Gly Val Thr Thr Gly
      115             120             125

Thr Pro Ile Leu Val Ile Val Pro Asn Thr Asp Gln Ile Gly Ser Asp
      130             135             140

His Arg Glu Ile Ala Asn Val Tyr Arg Pro Ser His Ala Asp Ala Thr
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Tyr Asp Phe Lys Tyr Gly Val Arg Ala Val Gln Gly Gly Gly Arg Ser
      165             170             175

Ser Gly Arg Lys Thr Val Gly Arg Val Ala Ala Gly Ala Leu Pro Lys
      180             185             190

Lys Ile Leu Lys Leu Lys Cys Gly Leu Glu Ile Leu Ser Phe Val Ser
      195             200             205

Lys Val His Gln Val Val Leu Pro Glu Asp Ala Val Asp Tyr Gly Ser
      210             215             220

Val Thr Leu Glu Gln Ile Glu Ser Asn Ile Val Arg Cys Pro Asp Pro
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Glu Tyr Ala Glu Lys Met Ile Asp Ala Ile Asp Arg Val Arg Val Arg
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Gly Asp Ser Val Gly Gly Val Ile Thr Cys Val Ala Arg Asn Val Pro

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| 260 | | | | | 265 | | | | | 270 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Gly | Leu | Gly | Ser | Pro | Val | Phe | Asp | Lys | Leu | Glu | Ser | Glu | Leu | Ala |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Lys | Ala | Met | Leu | Ser | Ile | Pro | Ala | Ser | Asn | Gly | Phe | Glu | Ile | Gly | Ser |
| | | 290 | | | | | 295 | | | | | 300 | | | |
| Gly | Phe | Ala | Gly | Thr | Asp | Leu | Thr | Gly | Ser | Glu | His | Asn | Asp | Glu | Phe |
| | | | | | | | 310 | | | | | 315 | | | 320 |
| Tyr | Met | Asp | Lys | Ala | Gly | Ser | Val | Arg | Thr | Arg | Thr | Asn | Arg | Ser | Gly |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Gly | Val | Gln | Gly | Gly | Ile | Ser | Asn | Val | Glu | Ile | Val | His | Phe | Lys | Val |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Ala | Phe | Lys | Pro | Thr | Pro | Ser | Ile | Gly | Val | Lys | Gln | Asn | Thr | Val | Ser |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Arg | Glu | Arg | Gln | Asn | Val | Glu | Leu | Leu | Ala | Arg | Gly | Arg | His | Asp | Pro |
| | | 370 | | | | | 375 | | | | | 380 | | | |
| Cys | Val | Ala | Pro | Arg | Ala | Val | Pro | Val | Val | Glu | Ser | Met | Ala | Ala | Leu |
| | | | | 390 | | | | | | | | 395 | | | 400 |
| Val | Leu | Met | Asp | Gln | Leu | Met | Ala | His | Val | Ala | Gln | Cys | Glu | Met | Phe |
| | | | | 405 | | | | | 410 | | | | | 415 | |
| Ala | Leu | Asn | Thr | Ala | Leu | Gln | Glu | Pro | Val | Gly | Ser | Phe | | | |
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 <212> DNA
 <213> Oryza sativa

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 ttagtcctca tggaccagct gatggcgcac attgctcaat gtgagatgtt tccactgaac 180
 cttgctctac aagaaccagt tggctctgcc agcagcgtac ctgcatttgc accagatcta 240
 aanngnccc ccctccccc ccccagctt gtttatcatc tatcatattt ctgggggttt 300
 ctaaggggtt cgcagttttg ccacaaagcc tgtatcctag tttatatctc gagttattgt 360
 acccaaggaa tccgtttatac agtgagcatg aagatagaaa tgcgttcacg cgtgttttgt 420
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<210> 10
 <211> 81
 <212> PRT
 <213> Oryza sativa

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 Pro Met Val Glu Ser Met Ala Ala Leu Val Leu Met Asp Gln Leu Met
 35 40 45
 Ala His Ile Ala Gln Cys Glu Met Phe Pro Leu Asn Leu Ala Leu Gln
 50 55 60
 Glu Pro Val Gly Ser Ala Ser Ser Val Pro Ala Phe Ala Pro Asp Leu
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 Ser

<210> 11
 <211> 966
 <212> DNA
 <213> Oryza sativa

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 cggagtcgcg cccggcttcc ctccgcttct ccgtcggccg ccgccgcgcc gctcgcctag 180
 aggtgaaggc gtctgcaaat gtatttgga actacttcca ggttgcaact tatggagagt 240
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 tacta 966

<210> 12
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 <213> Oryza sativa

<400> 12
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Leu Arg Phe Ser Val Gly Arg Arg Arg Ala Ala Arg Leu Glu Val Lys
 35 40 45
 Ala Ser Ala Asn Val Phe Gly Asn Tyr Phe Gln Val Ala Thr Tyr Gly
 50 55 60
 Glu Ser His Gly Gly Gly Val Gly Cys Val Ile Ser Gly Cys Pro Pro
 65 70 75 80
 Arg Ile Pro Leu Thr Glu Ala Asp Met Gln Val Glu Leu Asp Arg Arg
 85 90 95
 Arg Pro Gly Gln Ser Arg Ile Thr Thr Pro Arg Lys Glu Thr Asp Thr
 100 105 110
 Cys Lys Ile Leu Ser Gly Thr His Glu Gly Met Thr Thr Gly Thr Pro
 115 120 125
 Ile His Val Phe Val Pro Asn Thr Asp Gln Arg Gly Gly Asp Tyr Ser
 130 135 140
 Glu Met Ala Lys Ala Tyr Arg Pro Ser His Ala Asp Ala Thr Tyr Asp
 145 150 155 160
 Phe Lys Tyr Gly Val Arg Ala Val Gln Gly Gly Gly Arg Ser Ser Ala
 165 170 175
 Arg Glu Thr Ile Gly Arg Val Ala Ala Gly Ala Leu Ala Lys Lys Ile
 180 185 190
 Leu Lys Leu Lys Ser Gly Val Glu Ile Leu Ala Phe Val Ser Lys Val
 195 200 205
 His Gln Val Val Leu Pro Glu Asp Ala Val Asp Tyr Asp Thr Val Thr
 210 215 220
 Met Glu Gln Ile Glu Ser Asn Ile Val Arg Cys Pro Asp Pro Glu Tyr
 225 230 235 240
 Ala Gln Lys Met Ile Asp Ala Leu Asp Lys Val Arg Val Arg Gly Asp
 245 250 255
 Ser Ile Gly Gly Val Val Thr Cys Ile Ala Arg Asn Val Pro Arg Gly
 260 265 270
 Ile Gly Ser Pro Val Phe Asp Lys Leu Glu Ala Glu Leu Ala Lys Ala
 275 280 285
 Met Leu Ser Leu Pro Ala Ser Lys Gly Phe Glu Ile Gly Ser Gly Phe
 290 295 300

Val Phe Thr
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<210> 13
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 <212> DNA
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<220>
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<220>
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 <222> (436)..(437)
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<220>
 <221> unsure
 <222> (448)
 <223> n = A, C, G, or T

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 <222> (466)
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 <222> (473)
 <223> n = A, C, G, or T

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 <222> (479)
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 <222> (504)
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 <222> (522)
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 cccgcctacc tccgactctc actcgcgtcct cgtcttccca agagacttca catacaggcg 180
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 caagtggatc ttgacagaag gaggccaggt cagagccgaa ttacaactcc tagaaaggag 360
 actgatacat gtaaaatatt ttcaggagtt tccgaaggaa tcactactgg nactccaatt 420
 catgtactgt acccanntac tgatcaanga gggcatgact atagcnagat ggnagtacnt 480
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 g 541

<210> 14

<211> 168
 <212> PRT
 <213> Glycine max

<220>
 <221> UNSURE
 <222> (139)
 <223> Xaa = ANY AMINO ACID

<220>
 <221> UNSURE
 <222> (164)
 <223> Xaa = ANY AMINO ACID

<400> 14
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 Ala Phe Ala Ser Leu Asn Ser Asp Leu Gly Ser Leu Ser Pro Ala Tyr
 20 25 30
 Leu Arg Leu Ser Leu Arg Pro Arg Leu Pro Lys Arg Leu His Ile Gln
 35 40 45
 Ala Ala Gly Ser Thr Tyr Gly Asn His Phe Arg Val Thr Thr Tyr Gly
 50 55 60
 Glu Ser His Gly Gly Gly Val Gly Cys Val Ile Asp Gly Cys Pro Pro
 65 70 75 80
 Arg Leu Pro Leu Ser Glu Ala Asp Met Gln Val Asp Leu Asp Arg Arg
 85 90 95
 Arg Pro Gly Gln Ser Arg Ile Thr Thr Pro Arg Lys Glu Thr Asp Thr
 100 105 110
 Cys Lys Ile Phe Ser Gly Val Ser Glu Gly Ile Thr Thr Gly Thr Pro
 115 120 125
 Ile His Val Ser Val Pro Asn Thr Asp Gln Xaa Arg His Asp Tyr Ser
 130 135 140
 Glu Met Ala Leu Leu Ile Gly Leu His Ala Asn Ala Thr Tyr Asp Met
 145 150 155 160
 Lys Tyr Gly Xaa Arg Ser Val Lys
 165